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Molecular Transporter: Bacteriophage

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Commentary Article

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ABSTRACT

Enduring developments in constructing artificial DNA amalgamation and sequencing advances coupled with new methodology for genomic change and get together have opened the entryway for outfitting the force and contrasting characteristics of science for applications; such an assortment of procedures grow in the quality update. The present talk discuss the bacteriophage is a novel transporter of the molecular gene with the error gene.

Introduction

Gene therapy has pulled in extensive consideration in the previous couple of decades. One of its most encouraging applications is in the field of the repairing the error gene with normal. On the other hand, a few impediments need to be overcome before this possibly capable procedure can be clinically actualized. It is vital to recognize a systemically managed vector that can securely and viably convey restorative qualities to correctly focus on cells anyplace in the body [1-3].

Various late examinations have investigated tissue-focused on quality conveyance. Notwithstanding segregating target cells with a high level of exactness, this minimizes antagonistic symptoms, for example, cytotoxicity and resistant reaction by lessening the required vector load [4-6].

Inside the most recent 25 years, Bacteriophage have quickly ascended to as hereditary apparatus for an extensive variety of utilizations from basic cloning to genome designing due its genome size and high copy number [7].

Structure of Bacteriophage

Virus infected bacteria, or Bacteriophage, are pervasive creatures crossing altogether different natural specialties [8]. Despite the fact that genome examination neglects to show broad relationship among Bacteriophage, late auxiliary studies uncover a high level of likenesses. Most Bacteriophage presents an icosahedral proteinaceous head, which contains the nucleic corrosive, either deoxyribonucleic acid (DNA) or ribonucleic acid (RNA). Exemptions to this standard are couple of situations where a lipid envelope structures some piece of the head, and those different situations where the head exhibits a filamentous geometry [9-14].

Bacteriophages have made essential commitments to the field of sub-atomic science. In light of their basic structures and little number of qualities, bacteriophages are great exploratory frameworks for

hereditary building and investigating sub-atomic natural courses of action. In 1952, Hershey and Chase explained the hereditary capacity of DNA utilizing the T2 bacteriophage, which secured the establishments for in the long run affirming that DNA was the essential hereditary material. In acknowledgment of this major exploratory accomplishment, in 1969 Alfred Hershey and two different researchers imparted the Nobel Prize in Physiology or Medicine [15-17].

Use of Bacteriophage is picking up consideration as an option strategy for keeping the bacterial pollution. It is considered as a potential option biocontrol system to repress the pathogen. Harmful phages cause bacterial host cell lysis and capacity to control bacterial populaces as well as can be utilized as markers of bacterial sully in fecal examples and as a potential instrument for recognizing particular bacterial strains. A multivalent harmful Bacteriophage would be a decent determination for phage treatment as a result of its wide host range [18-20].

Bacteriophages as Potential Bioterrorism Agents or Tools

Transduction is one of the fundamental main impetuses of level quality move in microbial advancement, and it is likewise an imperative system for adjusting bacterial destructiveness, in light of the fact that lysogenic bacteriophages frequently convey harmfulness qualities [21-24]. A decent illustration of this is the *Vibrio cholerae* poison quality *ctxABT* that causes sustenance harming and is conveyed by its lysogenic bacteriophage.

As microbes with numerous medication resistance phenotypes get to be more regular, alongside better comprehension of drug resistance systems and anti-microbial capacity, it has gotten to be anything but difficult to get microscopic organisms that are increase drug-safe. Such microorganisms could be acquired through clinical gathering of medication safe microscopic organisms or by fake blend of different medication resistance qualities and harmfulness qualities [25-27].

Application of bacteriophages for bacterial tracing and typing could be of immense value when dealing with an unexpected bioterrorist attack or bacterial disease epidemic [28-31].

Bioterrorism or biothreat specialists are exceedingly irresistible and pathogenic microorganisms (microscopic organisms, infections, and parasites) and their poisons that can be utilized by people or gatherings of terrorists or as biowarfare operators in military operations [32-33].

The seriousness of these biothreat bacterial contaminations requires proficient biosurveillance and biodefense, including accessibility of a rich stockpile of exceptional techniques for fast discovery and recognizable proof of the microscopic organisms, strain portrayal, diagnostics, effective prophylaxis, and treatment of these diseases. Numerous bacterial infections (bacteriophages or phages) dynamic against *Y. pestis*, *B. anthracis* and *Brucella* species have been depicted [34-36]. Attributes and handy utilizations of such phages are the subject of this survey. Because of the absence of information on lytic phages of *F. tularensis* and exceptionally restricted data on viable significance of phages fit for lysing *B. pseudomallei* and *B. mallei*, this audit does not cover the writing on these microorganisms [37-40].

Bacteriophage-based Diagnostics

Routine bacteriophage lysis tests have been used for identification of biothreat bacteria, discrimination of relative species and differentiation of typical and atypical strains for more than 80 years [96,100,106]. In particular, the phage lysis assay is an essential part of *Y. pestis* identification and bacteriological diagnosis of plague [41-43].

Coming to environmental applications there are two critical fields in natural utilizations of bacteriophages: recognition of biothreat microscopic organisms in ecological examples and phage-based cleaning. Rather than PCR tests focusing on bacterial DNA, phage-construct strategies that depend with respect to phage proliferation distinguish just live bacterial cells. This is an unequivocal focal point when testing the movement of common foci of malady and for measurable purposes, for example, amid the examination of bioterrorist assaults [44].

Genome and size

The hereditary differing qualities of the bacteriophage populace are striking. All in all, the nucleotide arrangements of genomes got from phages with non-covering host goes seldom impart succession closeness, despite the fact that this may not be astonishing if the bacterial hosts are indirectly related. Since bacteriophages tainting a typical bacterial host are in hereditary contact with one another, it is not astounding that they now and again impart normal nucleotide successions [45].

Bacteriophage is a lytic phage that pollutes enterobacteria *Escherichia coli*. It is one of the best-depicted phage in the T5-like disease's assortment of the Siphoviridae Family. Sequenced phage genomes vary broadly in size from *Leuconostoc* phage L5 (2,435bp) to *Pseudomonas* phage 201phi2-1 (316,674b). In any case, there is not a uniform transport of genome sizes over this reach [46].

Of the 168 putative ORFs, 61 (36.3%) have been consigned limits according to their homology to known groupings. These are qualities primarily included in DNA replication and repair, nucleotide absorption framework, lysis, phage helper proteins and distinctive chemicals Fifteen (8.9%) are proteins that match to hypothetical proteins. Ninety-two (54.7%) are expected ORFs lacking similarity to any known proteins [47].

Treatments for muscular dystrophies remain a real test disregarding propelled procedures utilizing either cell or quality treatment. We here propose a joined methodology of cell and quality treatment. As quality conveyance vehicles with particular homing potential we have picked mesoangioblasts which are undeveloped cells with mesodermal potential.

Progressing remedial methodologies intend to enhance the indications of solid dystrophies of which Duchenne strong dystrophy (DMD) is among the most extreme. Procedures incorporate pharmaceutical, hereditary and cell restorative routines or mixes of these. So far serious examination has not possessed the capacity to ease this hereditary threatening muscle squandering ailment essentially because of the far reaching dispersion of skeletal muscle in the body [48].

The genetic contrasting characteristics of the bacteriophage people are magnificent. At the point when all is said in done, the nucleotide plans of genomes got from phages with non-covering host expands rarely bestow gathering closeness, notwithstanding the way that this may not be stunning if the bacterial hosts are in a roundabout way related. Since bacteriophages polluting an ordinary bacterial host are in inherited contact with each other, it is not stunning that they at times offer essential nucleotide plans. More than 30 phage genomes have been separated for *Pseudomonas* comprise of 33, *Staphylococcus* around 48 and *Mycobacterium* comprise of 50 hosts, and there are various situations where phages of a commonplace host acknowledge related progression [49].

From Normal Cloning Vectors to Molecular transporter

Different sorts or sorts of cloning vectors are utilized for the exchanging of nature of enthusiasm into the searched for tissue or cell. Subsequently vectors are utilized as vehicle to go on nature of vectors. A few sorts of vectors are being made in lab; every vector has got unmistakable sub-atomic properties and additionally cloning most extreme. A rate of the cases for vectors utilized as a bit of recombinant DNA progression are plasmids, phagemids, cosmids, shuttle vectors and essentially more for the cloning of the quality [50].

CONCLUSION

One reason that phages have been important in investigation is the effortlessness of synchronizing the lytic cycle in a masses of cells, so that the development of sub-nuclear events can be measured over the

whole society. Synchronization may be fulfilled either by simultaneous pollution or by the affectation of phage change in lysogenic cells. A second reason is the straightforwardness with which changes that impact specific periods of the cycle can be differentiated and separated. Phages are as vacillated fit as a fiddle as the diseases that pollute eukaryotes. A couple of formal portrayal arrangements have been proposed, yet none is extensively useful or known to reflect phylogeny. In light of the current writing and condition of the field, one can reason that Bacteriophage lytic routinely utilized for the recognition, recognizable proof and writing of the host microorganisms.

REFERENCES

1. Fan H, Tong Y. Potential Dual-Use of Bacteriophage Related Technologies in Bioterrorism and Biodefense. *J Bioterr Biodef.* 2012; 3:121
2. Patriarca A, Salutari P, Di Zacomo S. The Impact of Molecular Genetic in Acute Myeloid Leukemias. *J Blood Disorders Transf.* 2015; 6:252
3. Bhensdadia DV, Bhimani HD, Nathani NM, Rawal CM, Koringa PG, et al. Isolation, Molecular Characterization and Insight into the Genome Sequence of E. Coli Bacteriophage ADB-2 from Poultry Fecal Sample. *Next GeneratSequenc&Applic.* 2014; 1:101
4. Filippov AA, Sergueev KV, Nikolich MP. Bacteriophages against Biothreat Bacteria: Diagnostic, Environmental and Therapeutic Applications. *J Bioterr Biodef* 2013; S3:010
5. Nora Beatriz Calcaterra. How to Have Dual Lives: Proteins that Bind DNA and RNA. *MolBiol.* 2014; 3:e120
6. Dutt S, Tanha J, Evoy S, Singh A. Immobilization of P22 Bacteriophage Tailspike Protein on Si Surface for Optimized Salmonella Capture. *J Anal Bioanal Techniques.* 2013; S7:007
7. Singh PK. Vitamin E Analogs as Radiation Counter measures: Beyond the Antioxidant Activities. *Mol Biol.* 2014; 3:e116
8. Patsouris D, Jeschk MG. Stress Induced Insulin Resistance in Regards to Cellular Organelles, Inflammasome and Inflammation and Lipids. *MolBiol.* 2014; 3:e114.
9. Nallaseeth. Is There a Role for an Evolutionary Genetics Based Rational Health Policy In Global Biomedical, Health and Economic Policies? *.MolBiol.* 2014; 3:1000e118
10. Proost J, Deschuyteneer G, Santoro R, Overmeere QV, Soumillion P, et al. Filamentous Phages Displaying Multivalent Peptide Motives with Specific Affinity to Anodic Alumina Surfaces. *J Bioeng Biomed Sci.* 2015; 6:162
11. Takis. A Celebrating 30 Years since the Conception of the Human Genome Project (HGP): New Concepts Ahead-Molecular Biology Tools to Efficiently Modify the HG and/or Other Species-Genomes-Implications for Health and Disease. *MolBiol.* 2014; 3:e119
12. Bhowmick S, Tripathy S. A Tale of Effectors; Their Secretory Mechanisms and Computational Discovery in Pathogenic, Non-Pathogenic and Commensal Microbes. *Mol Biol.* 2014; 3:118
13. Remes AM, et al. Functional MRI in Patients with the C9ORF72 Expansion associate Frontotemporal Dementia. *MolBiol.* 2014; 3:117
14. Perepechaeva M, Kolosova N and Grishanova A. Altered mRNA Expression of “Ahr-Nrf2 Gene Batteries” in the Retinas of Senescence-Accelerated OXYS Rats during Development of AMD-Like Retinopathy. *J Mol Genet Med.* 2014; 8:105
15. Zilber-Rosenberg I, Rosenberg E. Prebiotics and Probiotics within the Framework of the Hologenome Concept. *J Microbial BiochemTechnol.* 2011; S1:001
16. Turton JF, Perry C, Hannah MJ. Isolation of Bacteriophage against Currently Circulating Strains of *Acinetobacterbaumannii*. *J Med Microb Diagn.* 2012; 1:109
17. Higgins PJ. PAI-1 Promoter-specific Oligonucleotide Decoys: Transcription Factor α Bates and Potential Utility as Wound Healing Therapeutics. *Cell Dev Biol,* 2014; 3:143
18. Agyare C, Ansah AO, Ossei PPS, Apenteng JA, Boakye YD. Wound Healing and Anti-Infective Properties of *Myrianthusarboreus* and *Alchorneacordifolia*. *Med chem.* 2014; 4:533-539
19. Collawn SS, Patel S. Adipose-Derived Stem Cells, their Secretome, and Wound Healing. *J Cell Sci Ther.* 2014; 5: 165
20. Wlian L, Carrijo-Carvalho LC, Andrade SA, Lourenço SV, Rodrigues CJ, et al. Wound Healing Effects of a Lipocalin-Derived Peptide. *J Clin Toxicol.* 2014; 4:187

21. Marjeta U. Chemo Proteomics, a Valuable Tool for Biomarker and Drug Discovery. *Mol Biol.* 2014; 3:e117
22. Kajhøj TQ, Duch M, Pedersen FS, Løvschall H, Füchtbauer EM. Test of Critical Steps towards a Combined Cell and Gene Therapy Approach for the Treatment of Duchenne Muscular Dystrophy. *J Mol Genet Med.* 2015; 9:160.
23. Altaner C. Prodrug Gene Therapy for Cancer Mediated by Mesenchymal Stem/Stromal Cells Engineered to Express Yeast Cytosine deaminase::Uracil phospho ribosyl transferase. *J Stem Cell Res Ther.* 2015; 5:264
24. Venkatachalam KV. Science Vision 2020 on Sulfur Metabolism: What is Needed and What can be Achieved?. *J Genet Syndr Gene Ther.* 2015; 6:e129
25. Peter Morcos N, Ulrika Andersson ME, Eric Adler D. Left Ventricular Non-Compaction: Current Controversy and New Insights. *J Genet Syndr Gene Ther.* 2015; 6:255.
26. Arancio W, Genovese SI, Pizzolanti G, Giordano C. Hutchinson Gilford Progeria Syndrome: A Therapeutic Approach via Adenoviral Delivery of CRISPR/cas Genome Editing System. *J Genet Syndr Gene Ther.* 2015; 6:256
27. Fumikazu Koyama, Kazuaki Uchimoto, Hisao Fujii, Hirofumi Hamada, Kazuo Ohashi, et al. Adenovirus-Mediated Bcl-Xl Gene Therapy Combined with Pronase Treatment Protects the Small Intestine from Radiation-Induced Enteritis in Mouse Model. *J Genet Syndr Gene Ther.* 2014; 5:239
28. Avina Fierro JA and Hernandez Avina DA. A Case of Complete Cutaneous Syndactyly of the Toes with Non-Syndromic Phenotype. *J Genet Syndr Gene Ther.* 2014; 5:240
29. Rao GHR, Gandhi PG, Sharma V. Clinical Complications of Type- 2 Diabetes Mellitus in South Asian and Chinese Populations: An Overview. *J Diabetes Metab.* 2014; 5:420
30. Kajhøj TQ, Duch M, Pedersen FS, Løvschall H, Füchtbauer EM. Test of Critical Steps towards a Combined Cell and Gene Therapy Approach for the Treatment of Duchenne Muscular Dystrophy. *J Mol Genet Med.* 2015; 9:160
31. Ahmad I, Shahin R. Autologous Gene Therapy - A Proactive Approach to Cancer Critique. *J Cancer Sci.* 2014; Ther2014 R1:001
32. Riccobono D, François S, Valente M, Forcheron F, Drouet M. Advances in Stem Cell Therapy: Specific Applications in the Treatment of Cutaneous Radiation Syndrome. *J Stem Cell Res Ther.* 2014; 4:186
33. Zhang Y, Yao X. The Recent Advances and Future Perspectives of Personalized Medicine. *J Genet Syndr Gene Ther.* 2014; 5:e125
34. Berumen LC, García AG. 5-Ht5a Receptors during Ontogeny. *J Genet Syndr Gene Ther.* 2014; 5:e126
35. Berkinbayev S, Rysuly M, Mussayev A, Blum K, Baitasova N, et al. Apolipoprotein Gene Polymorphisms (APOB, APOC111, APOE) in the Development of Coronary Heart Disease in Ethnic Groups of Kazakhstan. *J Genet Syndr Gene Ther.* 2014; 5:216
36. Chang S, Gao L, Li Z, Ni L, Tong D, et al. Luciferase Expression is more Accurate than GFP to Assess Mirnas-Relevant Oncogenesis in vivo Live Imaging Study. *Cell Mol Biol.* 2014; 60: 104
37. Wegmann RJ. The Robberies and the Embezzlements Made to my Journal, *The Cellular and Molecular Biology@.* *Cell Mol Biol.* 2014; 60: 101
38. Ricard J. Emergence of Information and the Origins of Life a Tentative Physical Model. *Curr Synthetic Sys Biol.* 2014; 2:109
39. Yamane T, Sugimoto N, Maita H, Watanabe K, Takahashi-Niki K, et al. Identification of DJ-1-Associated Regions on Human Genes from SH-SY5Y Cells using Chromatin Immunoprecipitation Sequence Technique. *Mol Biol.* 2013; 3:115
40. Oliveira CAF. Preventing Pathogenic Bacteria in Milk and in Dairy Farms: The Usefulness of Molecular Biology Tools. *Adv Dairy Res.* 2013; 2:e105
41. Lakshmipathy U, Fontes A, Poderycki M, Chesnut JD. An Alternate Method for Efficient Delivery of Catalyzing Enzymes. *Mol Biol.* 2013; 2:112
42. Gopal P, Ragnath C, Vyas V, Shanmugam M, Ramasubbu N. Pub: Probing the Interaction of Human Salivary Alpha-Amylase and Amylase Binding Protein A (AbpA) of *Streptococcus gordonii*. *Mol Biol.* 2013; 2:111
43. Zhang K. Mass Spectrometry in Epigenetic Studies on Disease: Current Progress, Limitation, and Prospective. *Mol Biol.* 2013; 2:e109.
44. Zahur M, Asif AR. Clinical, Cellular & Molecular Biology of Autoimmune Disorders – Introduction. *J Clin Cell Immunol.* 2013; S10:e001.

45. Choene M, Motadi LR. Anti-Proliferative Effects of the Methanolic Extract of *Kedrostisfoetidissima* in Breast Cancer Cell Lines. *Mol Biol.* 2012; 1:107
46. Agostino AD, Gatta AL, Busico T, Rosa MD, Schiraldi C. Semiinterpenetrated Hydrogels Composed of PVA and Hyaluronan or Chondroitin Sulphate: Chemico-Physical and Biological Characterization. *J Biotechnol Biomater.* 2012; 2:140
47. Huang J, Wang L, Jiang M, Lin H, Qi L, et al. Parathyroid Hormone- Like Hormone (PTHrH) Feedback Mitosis to Downstream DNA Replication Coupling Postreplication Repair-Induced Cell Proliferation Network in No-Tumor Hepatitis/Cirrhotic Tissues (HBV or HCV Infection) by Systems-Theoretical Analysis. *Mol Biol.* 2012; 1:106
48. Qi L, Wang L, Jiang M, Huang J, Lin H. Cytosolic Iron-Sulfur Protein Assembly 1 (CIAO1) Downstream Activation of Phospholipase A2 and Hormone- Mediated Signaling-Induced Cell Death Network in Human Hepatocellular Carcinoma (HCC) by Systems-Theoretical Analysis. *Mol Biol.* 2012; 1:105
49. Lakshmipathy U, Fontes A, Poderycki M, Chesnut JD. An Alternate Method for Efficient Delivery of Catalyzing Enzymes. *Mol Biol.* 2013; 2:112
50. Perfilyeva Y, Ostapchuk Y, Cetin EA, Yilmaz A, Deniz G, et al. Hyaluronan-Binding T Regulatory Cells in Peripheral Blood of Breast Cancer Patients. *J Clin Cell Immunol.* 2015; 6:286